RAW SEQUENCE LISTING
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Application Serial Number: _____09 / 927,811B_____

Source: _____O IPE_____

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Revised 01/29/2002

Does Not Comply
Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/927,811B

DATE: 02/12/2003
TIME: 12:29:54

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF4\02122003\I927811B.raw

_Please re-submit in the English language.

```
   3 <110> APPLICANT: RheinBiotech Gesellschaft fur neue biotechnologische Prozesse
   4        und Produkte mbH
   6        Romano, Ivano
   8        Gellissen, Gerd
  10        DeVergilio, Claudio
  12 <120> TITLE OF INVENTION: Heat-Inducible Promoter
  14 <130> FILE REFERENCE: PCT1106-01966
  16 <140> CURRENT APPLICATION NUMBER: 09/927,811B
C--> 18 <141> CURRENT FILING DATE: 2003-02-07
  20 <150> PRIOR APPLICATION NUMBER: PCT/EP00/01144
  22 <151> PRIOR FILING DATE: 2000-02-11
E--> 24 <160> NUMBER OF SEQ ID NOS: (27) - found 28
  26 <170> SOFTWARE: PatentIn version 3.1
```

ERRORED SEQUENCES

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 601 <210> SEQ ID NO: 28
 602 <211> LENGTH: 15
 603 <212> TYPE: DNA
 604 <213> ORGANISM: Saccharomyces cerevisiae
W--> 605 <220> FEATURE:
 606 <223> OTHER INFORMATION: Heat shock element
 608 <400> SEQUENCE: 28
 609 ggaacagaac aatcg                                                    15
```

delete

```
W--> 611 (continued).(continued).(continued).(continued).(continued).(continued).(continued).
(con
W--> 612 tinued).(continued).(continued).(continued).(continued).(continued).(continued).
(continu
W--> 613 ed).(continued).(continued).(continued).(continued).(continued).(continued).
(continued).(
W--> 614 continued).(continued).(continued).(continued).(continued).(continued).(continued).
(cont
W--> 615 inued).(continued).(continued).(continued).(continued).(continued).(continued).
(continue
W--> 616 d).(continued).(continued).(continued).(continued).(continued).(continued).
(continued).(c
W--> 617 ontinued).(continued).(continued).(continued).(continued).(continued).(continued).
(conti
W--> 618 nued).(continued).(continued).(continued).(continued).(continued).(continued).
(continued
W--> 619 ).(continued).(continued).(continued).(continued).(continued).(continued).
(continued).(co
W--> 620 ntinued).(continued).(continued).(continued).(continued).(continued).(continued).
```

·(contin
W--> 621 ued) . (continued) . (continued) . (continued) . (continued) . (continued) . (continued) .
 (continued)
W--> 622 . (continued) . (continued) . (continued) . (continued) .
W--> 623 (8)
E--> 625 (1) -delete

} delete

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 02/12/2003
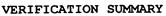PATENT APPLICATION: **US/09/927,811B**   TIME: 12:29:56

Input Set : **A:\PTO.VSK.txt**
Output Set: **N:\CRF4\02122003\I927811B.raw**

<u>Invalid Line Length:</u>
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:28; Line(s) 611,612,613,614,615,616,617,618,619,620,621

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/927,811B**

DATE: 02/12/2003
TIME: 12:29:56

Input Set : **A:\PTO.VSK.txt**
Output Set: **N:\CRF4\02122003\I927811B.raw**

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L:18  M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:71  M:283 W: Missing Blank Line separator, <220> field identifier
L:77  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:86  M:283 W: Missing Blank Line separator, <220> field identifier
L:92  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:109 M:283 W: Missing Blank Line separator, <220> field identifier
L:385 M:283 W: Missing Blank Line separator, <220> field identifier
L:389 M:283 W: Missing Blank Line separator, <220> field identifier
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:402 M:283 W: Missing Blank Line separator, <220> field identifier
L:404 M:283 W: Missing Blank Line separator, <220> field identifier
L:416 M:283 W: Missing Blank Line separator, <220> field identifier
L:430 M:283 W: Missing Blank Line separator, <400> field identifier
L:440 M:283 W: Missing Blank Line separator, <400> field identifier
L:447 M:283 W: Missing Blank Line separator, <220> field identifier
L:484 M:283 W: Missing Blank Line separator, <400> field identifier
L:494 M:283 W: Missing Blank Line separator, <400> field identifier
L:552 M:283 W: Missing Blank Line separator, <400> field identifier
L:563 M:283 W: Missing Blank Line separator, <400> field identifier
L:585 M:283 W: Missing Blank Line separator, <400> field identifier
L:605 M:283 W: Missing Blank Line separator, <220> field identifier
L:611 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:612 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:612 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:613 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:613 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:614 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:614 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:615 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:615 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:616 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:616 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:617 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:617 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:618 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:618 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:619 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:619 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:620 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:620 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:621 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:621 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:622 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:622 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:623 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:625 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:24  M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (27) Counted (28)
```